

Applicants: Zhang-Bao Xu, William S. Somers, Tatos N. Akopian,  
Chu-Lai Hsiao, and Raymond Unwalla  
Serial No.: 09/903,876  
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view of the above-noted amendments and these remarks, applicants respectfully submit that they have complied with the Sequence Rules. Accordingly, entry of the Sequence Listing is respectfully requested.

No fee is deemed necessary in connection with the filing of this Amendment. If any fee is required to preserve the pendency of the application, authorization is hereby given to charge the amount of any such fee to Deposit Account No. 01-1785.

Respectfully submitted,

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37174/9

SCHEDULE A

REDLINED VERSION

Please replace paragraph 13, at page 6, as follows:

Figure 1 provides a sequence alignment of ER- $\alpha$  (SEQ ID NO:1) with ER- $\beta$  (SEQ ID NO:2) covering the ordered extent of ER- $\beta$ . The numbering scheme used was chosen to be consistent with ER- $\alpha$ , such that the first ordered ER- $\beta$  residue, 311 is residue 263 in the full length protein. Residues in helix 12 are underlined. The (\*) symbols indicate the altered binding site residues.